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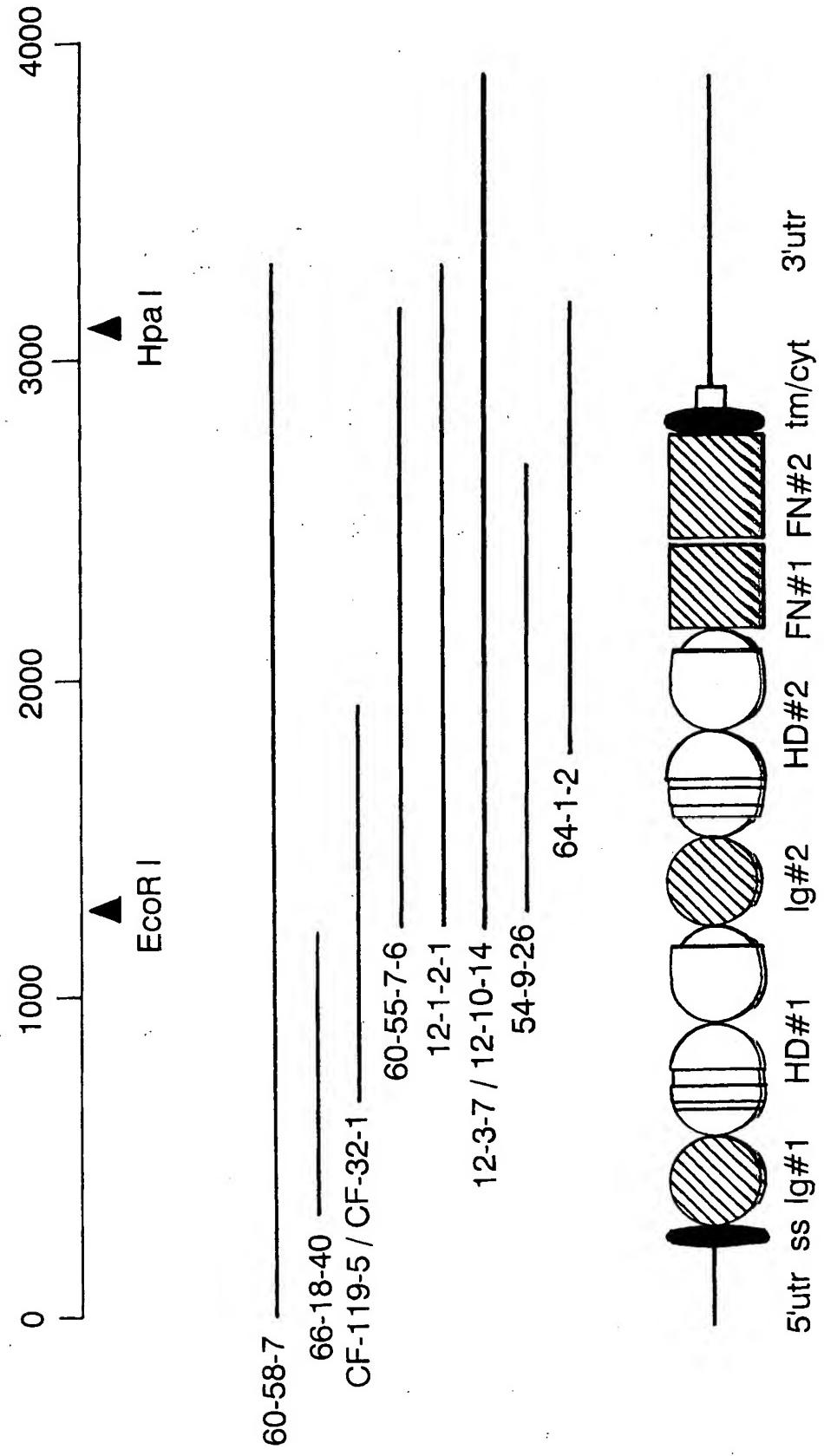


Figure 1



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-255	cgaaattcgcggggcgc
-240	gtcgacccggccaggctcgggagacatgggggggttaaaggctcgtnattatcc
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-120	tttggcagttcacccctgaccatcttggaaaataagttatctgtatctgtatgttt
-60	acttctccctcaccaacggagaacaaatgtggcaaaatgtgtacttctgtatgttt
1	ATGATTGTCAAAATTCTGTGGTTACATTGGAAATTATTATGTGATAACT
1	M I C Q K F C V V L L H W E F I Y V I T
61	GCGTTAACTTAATTCTGCATATCCAATTAAAGTTAAGTTGGAGATTCTGCCTTGCTTGCATGCCACCA
21	A F N L S Y P I T P W R F K L S C M P P
121	AATTCAACCTATGACTTACTTCCTTTGCCTGGACTCTCAAAGAATACTTCAAATTCCG
41	N S T Y D Y F L H P A G L S K N T S N S

**Figure 2A**



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A - - - - - A

181 AATGGACATTATGAGACAGCTGTTAACCTAAAGTTAACCAAGTGGTACTCACTTTCT  
61 N G H Y E T A V E P K F N S S G T H F S

241 AACTTATCCAAACTTCCACTGTTGCCTTCGGAGGTGAGCAAGATAAGAAACTGCTCC  
81 N L S K T T F H C C F R Q D R N C S

301 TTATGGCAGACAAACATTGGAAAGGACATTTGGTTCAACAGTAAAATTCTTTAGTTT  
101 L C A D N I E G R T F V S T V N S L V F

361 CAACAAATAGATgCAAACATGGAACATAACAGTGCTGGCTAAAAGGAGACTAAATTATTTC  
121 Q Q I D A N W N I Q C W L K G D L K L F

421 ATCTGTTATGGAGTCATTTAACGAAATTCAACTAAACTATAAGGTCCAT  
141 I C Y V E S L F K N L F R N Y N Y K V H

B - - - - - B

## Figure 2B



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B

481 CTTTTATATGGTCTGCCTGAAAGATTAGAAGATTACCTCTGGTTCCCCAAAAGGCCAGT  
161 L L Y V L P E V L E D S P L V P Q K G S

541 TTTAGATGGTCACTGCAATTGCAGTGTCAATGTCATGAAATGTTGTCAAATGTCATGCTGTG  
181 F Q M V H C N C S V H E C C E C L V P V

601 CCAACGCCAAACTAACGACACTCCCTATGTGTTGAAAATCACATCTGGTGGAGTA  
201 P T A K L N D T L L M C L K I T S G G V

661 ATTTTCCrGTCAACCTCTAAATGTCAGTTCAAGCCATAATGGTGAAGCCTGATCCACCA  
221 t F X S P L M S V Q P I N M V K P D P . P

721 TAGGTTGCATATGAAATCACAGATGATGGTAATTAAAGATTTCTGGTCCAGCCCCA  
241 L G L H M E I T D D G N L K I S W S S P

C — - C

Figure 2C



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C - - - - - C

781 CCATTGGTACCTCCACTTCAATTCAAGTGAATAATTCAAGGAAATTCTAGAGAATTACAAACAGTT  
261 P L V P F P L Q Y Q V K Y S E N S T T V

841 ATCAGAGAAGCTGACAAGATTGTCTCAGCTACATCCCTGCTAGTAGACAGTATACTTCCCT  
281 I R E A D K I V S A T S L L V D S I L P

901 GGGTCTTCGTTAGGGTTCAAGGTGAGGGCAAGAGACTGGATGCCAGGAATCTGGAGT  
301 G S S Y E V Q V R G K R L D G P G I W S

961 GACTGGAGTACTCCCTCGTGTCTTACCAAGATGTCAATACTTCCACCTAAAT  
321 D W S T P R V F T T Q D V I Y F P P K I

1021 CTGACAAAGTGGTCTAATGTTCTTTCACTGCATCTAAAGAAGGAAACAAAGAT  
341 L T S V G S N V S F H C I Y K K E N K I

D - - - - - D

## Figure 2D



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D ----- D

1081 GTTCCCTCAAAAGAGATTGTTGGATGAATTAGCTGAGAAAATTCCCTCAAAGCCAG  
361 V P S K E I V W W M N L A E K I P Q S Q

1141 TATGATGTTGAGTGCATGTTAGCAAAGTTACTTTCAATCTGAATGAAACCAA  
381 Y D V V S D H V S K V T F N L N E T K

1201 CCTCGAGAAAGTTACCTATGATGCAGTGCTGCAATGAAACATGCCATCAT  
401 P R G K F T Y D A V Y C C N E H E C H H

1261 CGCTATGCTGAATTATGATTGATGTCAATATCAATCTCATGTGAAACTGATGG  
421 R Y A E L Y V I D V N I S C E T D G

1321 TACTTAACCTAAATGACTTGCAGATGGTCAACCAGTACATCCAGTCACTTGGAAAGC  
441 Y L T K M T C R W S T S T I Q S L A E S

E ----- E

## Figure 2E



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E ----- E

1381 ACTTTGCAATTGAGGTATCATAGGAGCCAGCCTTACTGTTCTGATATTCCATCTATTCA  
461 T L Q L R Y H R S S L Y C S D I P S I H

1441 CCCATATCTGAGCCAAAGATTGCATATTGCAGAGTGATGGTTTATGAATGCATTTC  
481 P I S E P K D C Y L Q S D G F Y E C T F

1501 CAGCCAATCTTCCTATTATCTGGCTACACAAATGTGGATTAGGATCAAATCACTCTAGGT  
501 Q P I F L L S G Y T M W I R I N H S L G

1561 TCACTTGACTCTCCACCAACATGTGTCTCCTGATTCTGGTAAGCCACTGCCCTCCA  
521 S L D S P P T C V L P D S V K P L P P

1621 TCCAGTGTGAAAGCAGAAATTACTATAAACATTGGATTATTGAAATACTTGGGAAAG  
541 S S V K A E I T I N I G L L K I S W E K

F ----- F

## Figure 2F



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F ----- F

1681 CCAGTCTTTCCAGAGAAATAACCTTCAATTCCAGATTTCGCTATGGTTAACGTGGAAAAGAA  
561 P V F P E N N L Q F Q I R Y G L S G K E

1741 GTACAATGGAAGATGTATGAGGTTATGATCCAAACCACAAATCTGTCACTCCCAGTT  
581 V Q W K M Y E V D P K P K S V S L P V

1801 CCAGACTTGTGCAGTCTATGCTGTTCAAGGTCTTAAGAGGCTAGATGGACTGGGA  
601 P D L C A V Y A V Q V R F K R L D G L G

1861 TATTGGAGTAATTGGAGCAATCCAGCCTACACAGTTGATGGATATAAAGTTCTATG  
621 Y W S N W S N P A Y T V M D I K V P M

1921 AGAGGACCTGAATTGGAGATAATTAAATTGAGATACTATGAAAAAGGAGAAAAATGTC  
641 R G P E F W R I I N G D T M K K E K N V

G ----- G

## Figure 2G



G ----- G

1981 ACTTTACTTGGAAAGCCCCCTGATGAAAATGACTCATTGGCAGTGTTCAGAGATAATGTC  
661 T L L W K P L M K N D S L C S V Q R Y V

2041 ATAAACCATACATTTCCATSCAATGGAACATGGTCAGAAGATGTGGAAATCACACGAAA  
681 I N H H T S X N G T W S E D V G N H T K

2101 TTCACTTCTGTGGACAGAACATACTGTTACGGTTCTGGCCATCAATCAATT  
701 F T F L W T E Q A H T V T V L A I N S I T

2161 GGTGCTTCTGTTgCaAAATTAAATTAACTCTTTCATGGCCATAGGCAAAGTAATACT  
721 G A S V A N F N L T F S W P M S K V N I

2221 GTGCAGTCACTCAGTGCTTATCCTTAAACAGCAGTTGTGATGTTCCCTGGATACTA  
741 V Q S L S A Y F L N S S C V I V S W I L

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H ----- H

## Figure 2H



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H ----- H

2281 TCACCCAGT GATTACAAGCTTAATGTTATTGAGTGGAAAATCTTAATGAAGAT  
761 S P S D Y K L M Y F I I E W K N L N E D

2341 GGTGAAATAAAATGGCTTAGAATCTCTCATCTGTTAAGAAGTATTATCCATGATCAT  
781 G E I K W L R I S S V K Y T H D H

2401 TTTATCCCCATTGAGAACGTTCACTCAGTCAGTCTTACCCAATATTATGGAAAGGAGTGGGA  
801 F I P I E K Y Q F S L Y P I F M E G V G

2461 AACCAAAGATAATTAAATAGTTCAACTCAAGATGATATTGAAAAACACCAGAGTGTGCA  
821 K P K I I N S F T Q D D I E K H Q S D A

2521 GGTTTATGTAATTGTGCCAGTAATTTCCTCCATCTTGCCTTGGAAACATTA  
841 G L Y V I V P V I S S I L L G T L

I ----- I

## Figure 2!



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TTAATATCACACCAAGAACATTGGAAAGCTATTGGGAAGATGTCGGAACCCCCAAGAAT  
861 L I S H Q R M K L F W E D V P N P K N

TGTTCCCTGGGCACAGGACTTAATTTCAGAAAGAGAACGGACATCTTtgaagtctaatac  
881 C S W A Q G L N F Q K R T D I L \*

2701 atgatcactacagatgtccaaatgtccaaaccttccaaacacagtctatagagttatagaag  
3761 attttacatttgaaaggggaggcaaatctaaaaaaaattcaggttgaacttctgagag  
2821 ttaacatatggattatgtgatttqaacttaaaaatagatgtcattaaacccaagt  
2881 ttcatcataactcaggtaaacactaacactaaataaaagtttagatgttcaaatt  
2941 ttcatcataactaaaggaccggaaaactaaacagttataaggaccaggattttgttaattc  
3001 ttttaataccgacaacgacagtaatgtatagataatttacagtttatcacatctg  
3061 ttaggacattaaatccactttagatgttgcacgtttagactgtttatcgaaaattttatgt  
3121 tactaataattcatacccttagtcactttataaaatcaaaaatacaggtttgaaaaa

J. - - - - - J

**Figure 2J**



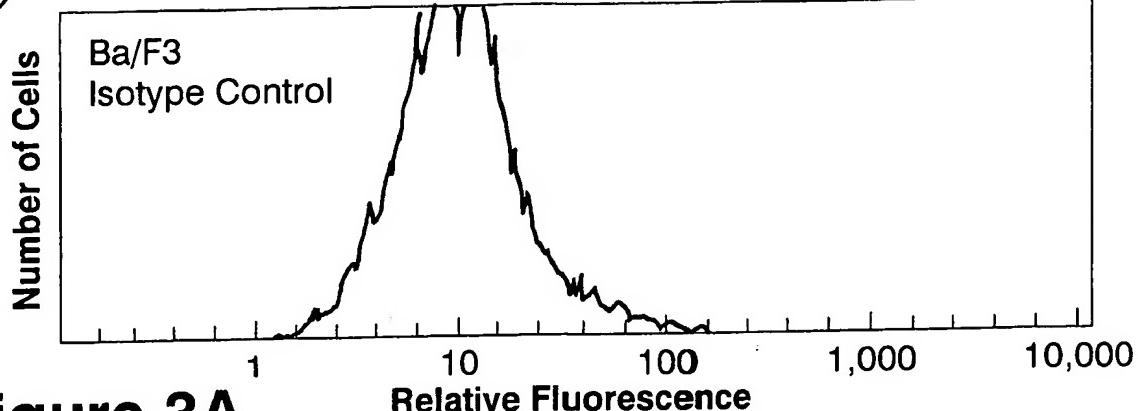
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3181 ggttaaatctaaggaaatatctgtgcagtggatttttagtcggataaggcccacaaagaaa  
3241 actttagaggaccgtaaaaacatagattgaaacaagtttagacccttaaagtcaaaaggtt  
3301 attaggaactttaccgaaattcactattgaaggccaaagtcaattttcccttgggctcaac  
3361 acaaacacgacgggtgtccctgtcacccctcaatgtcaagtagtatagtccctactggatgtatg  
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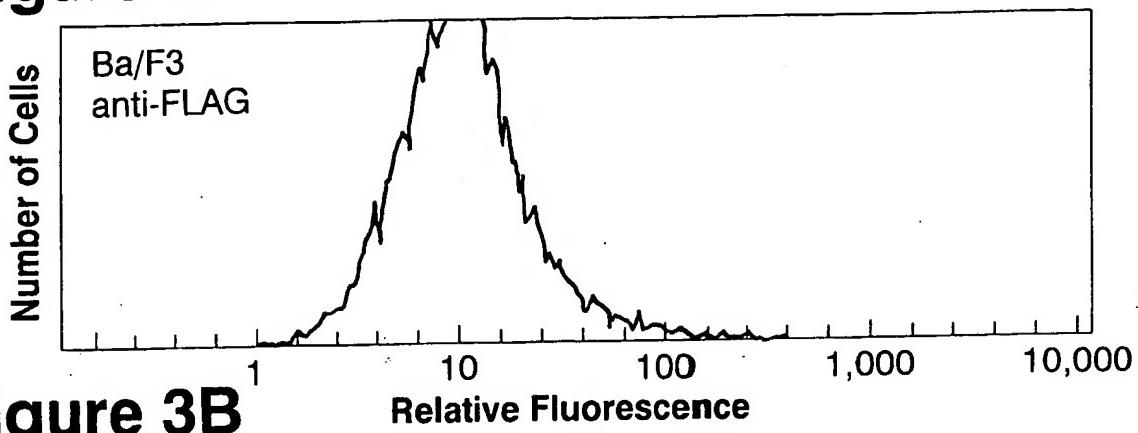
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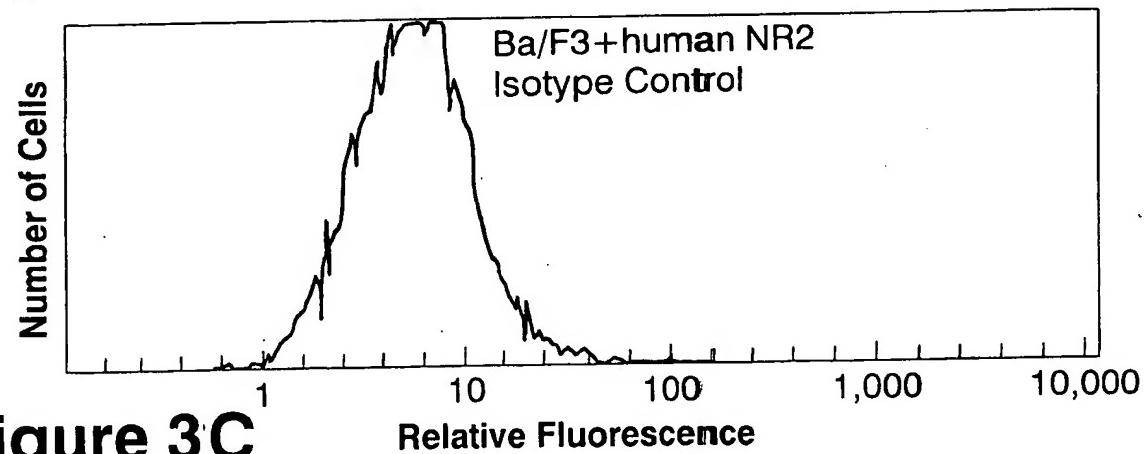
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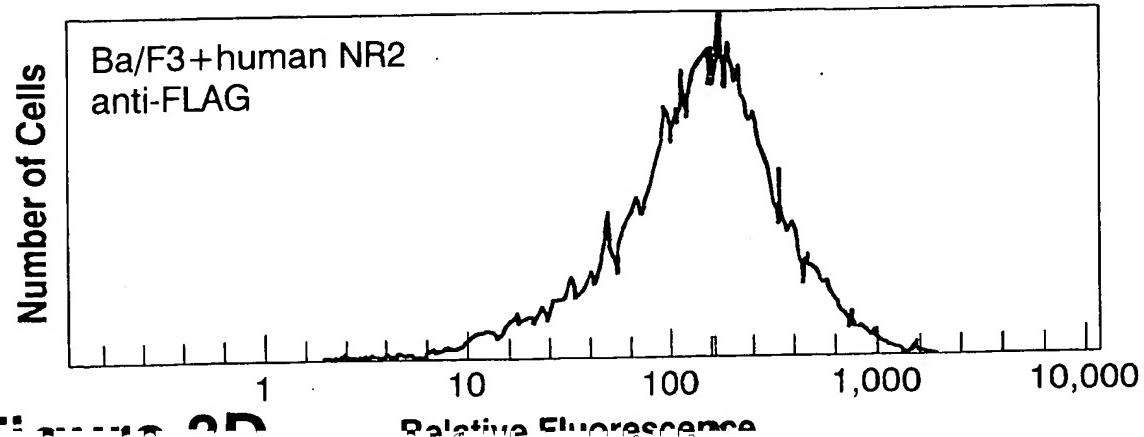
**Figure 3A**



**Figure 3B**



**Figure 3C**

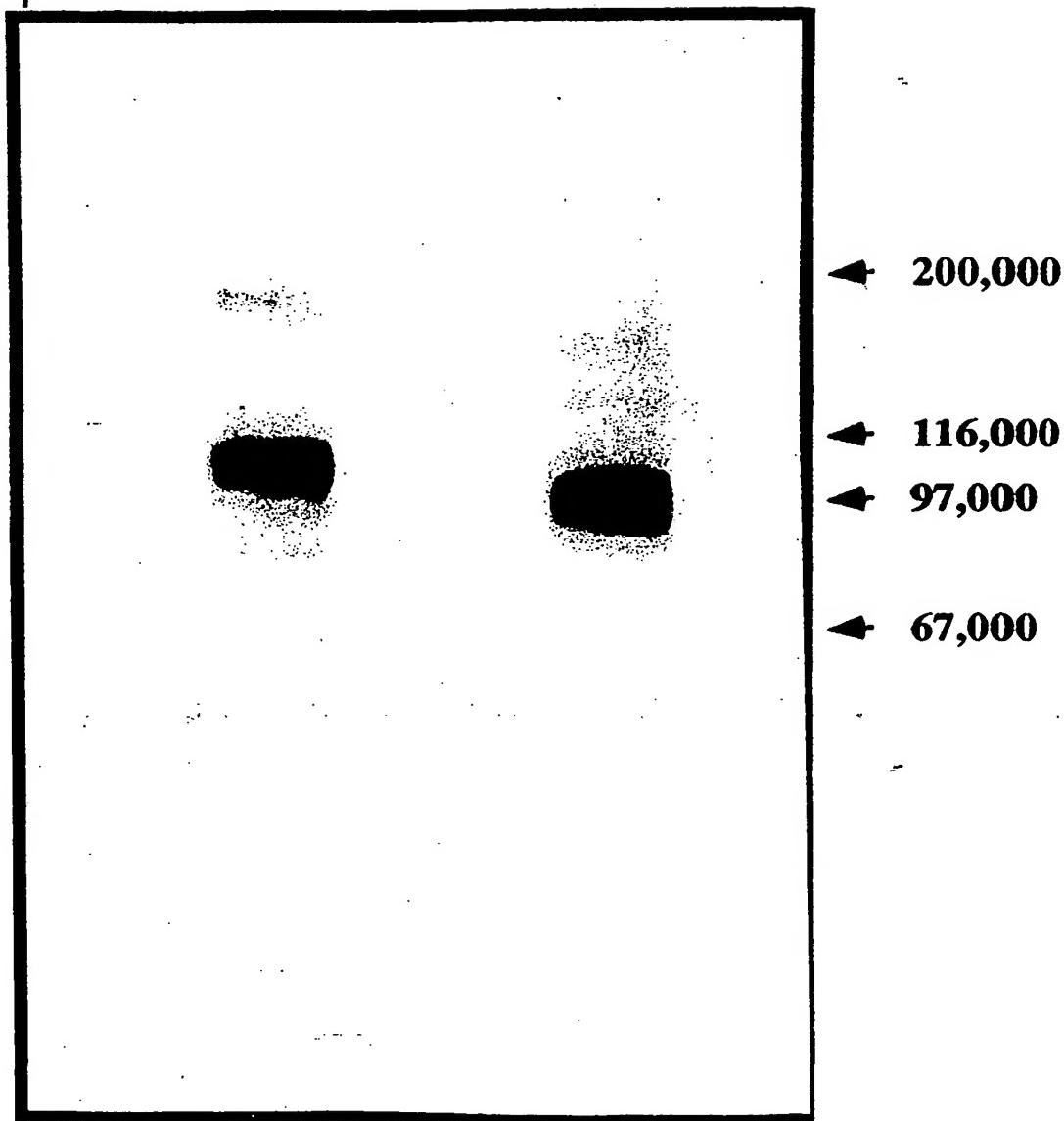


**Figure 3D**



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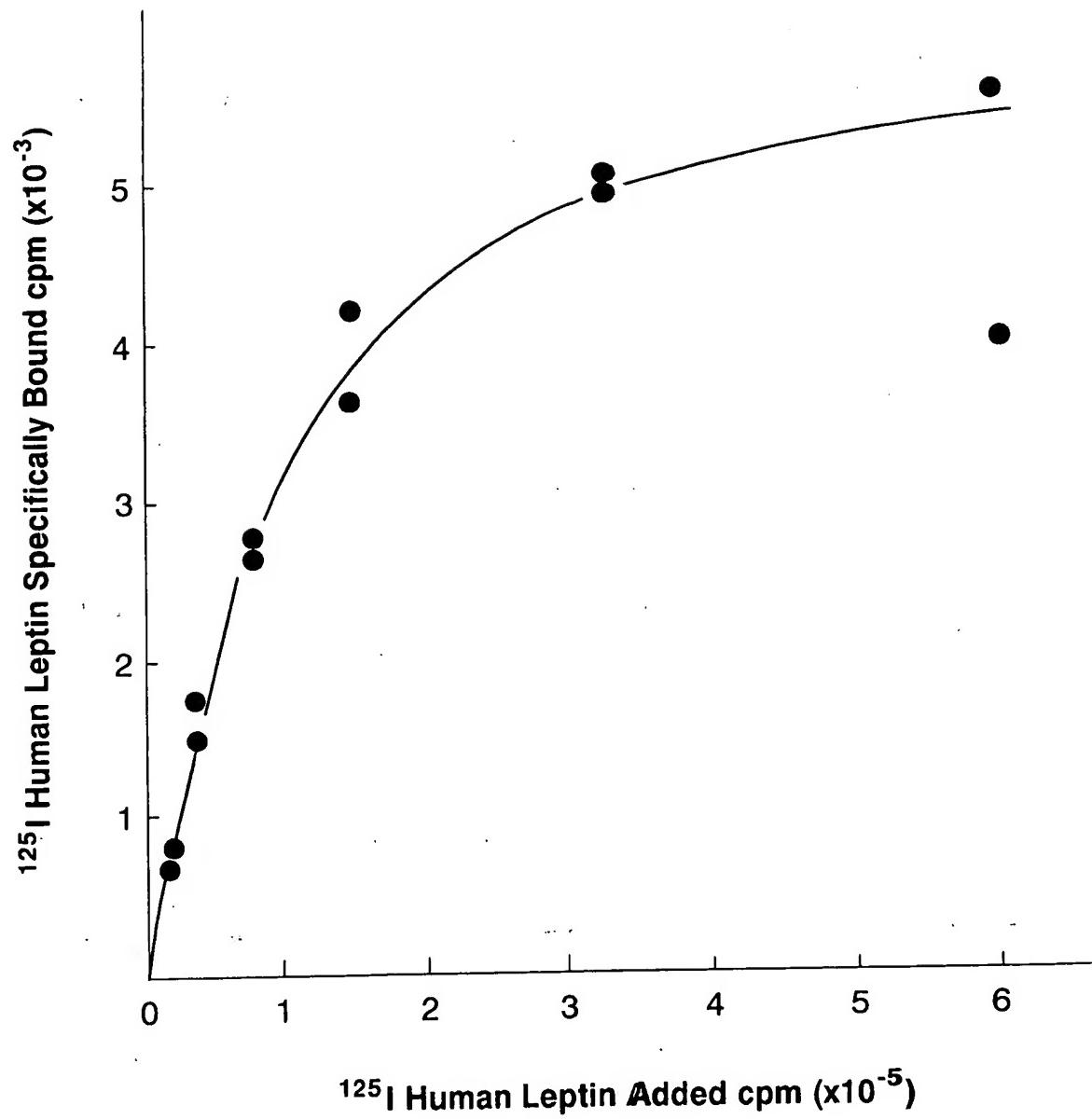
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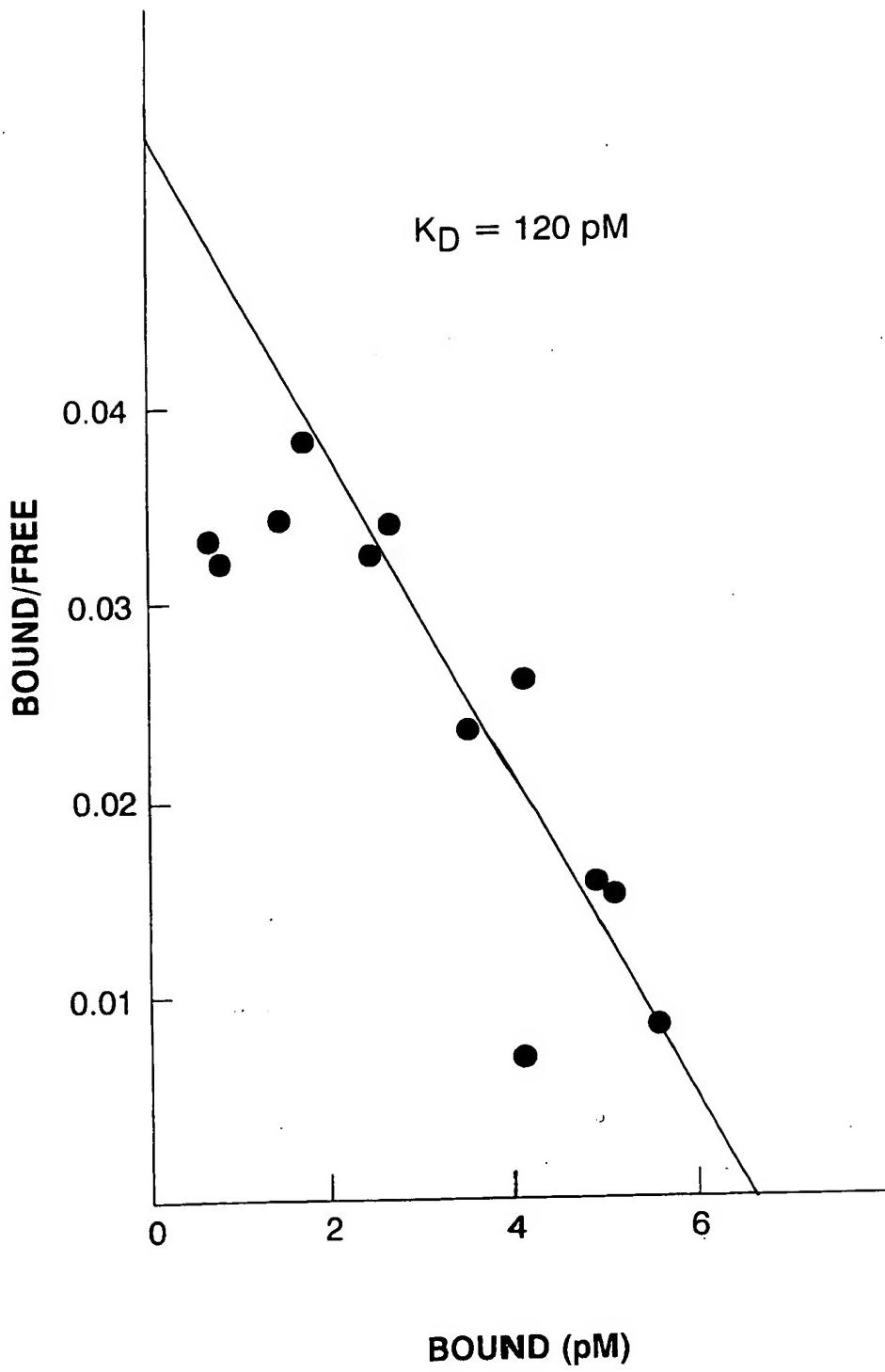
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**Figure 5A**



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**Figure 5B**



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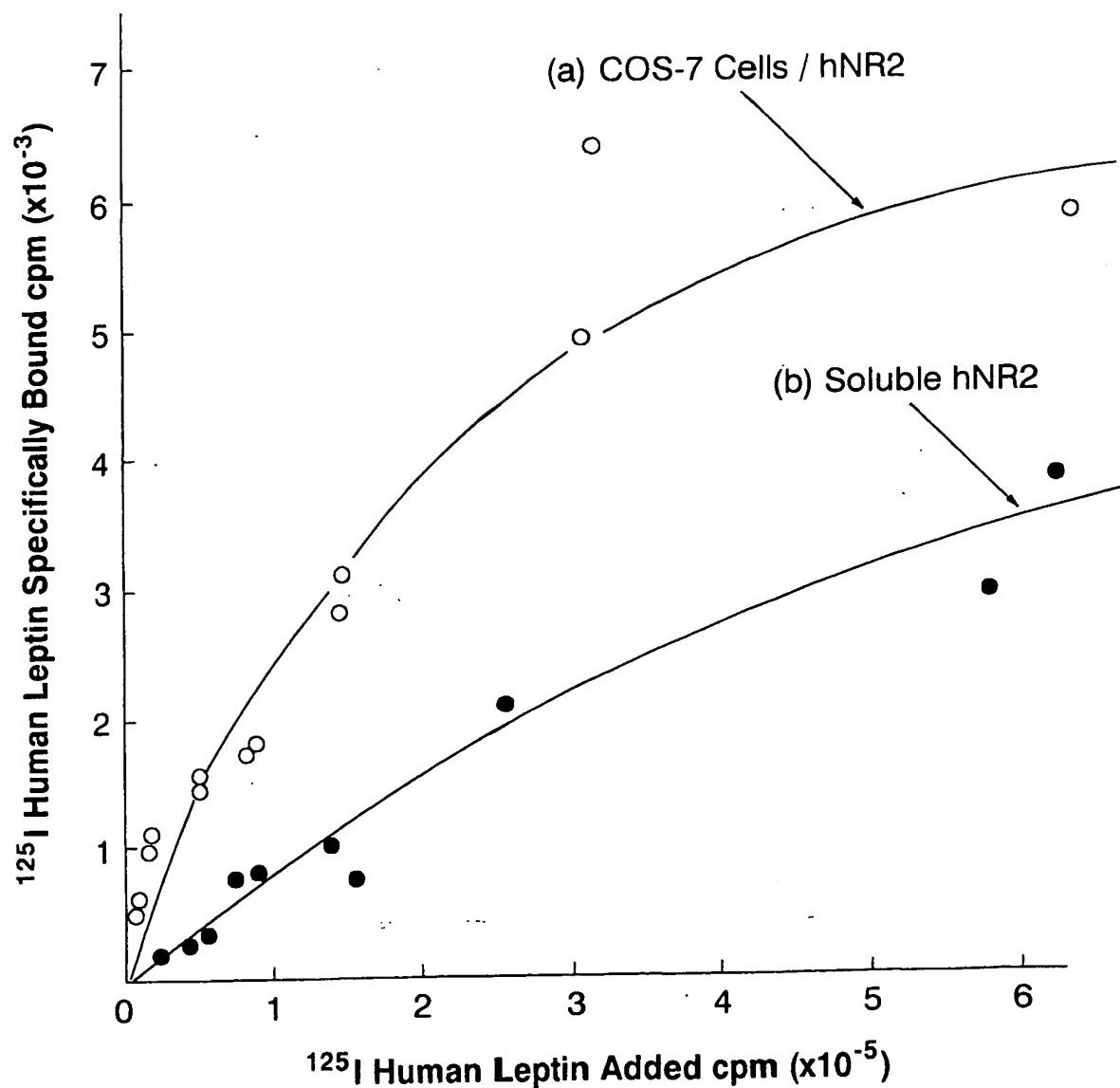
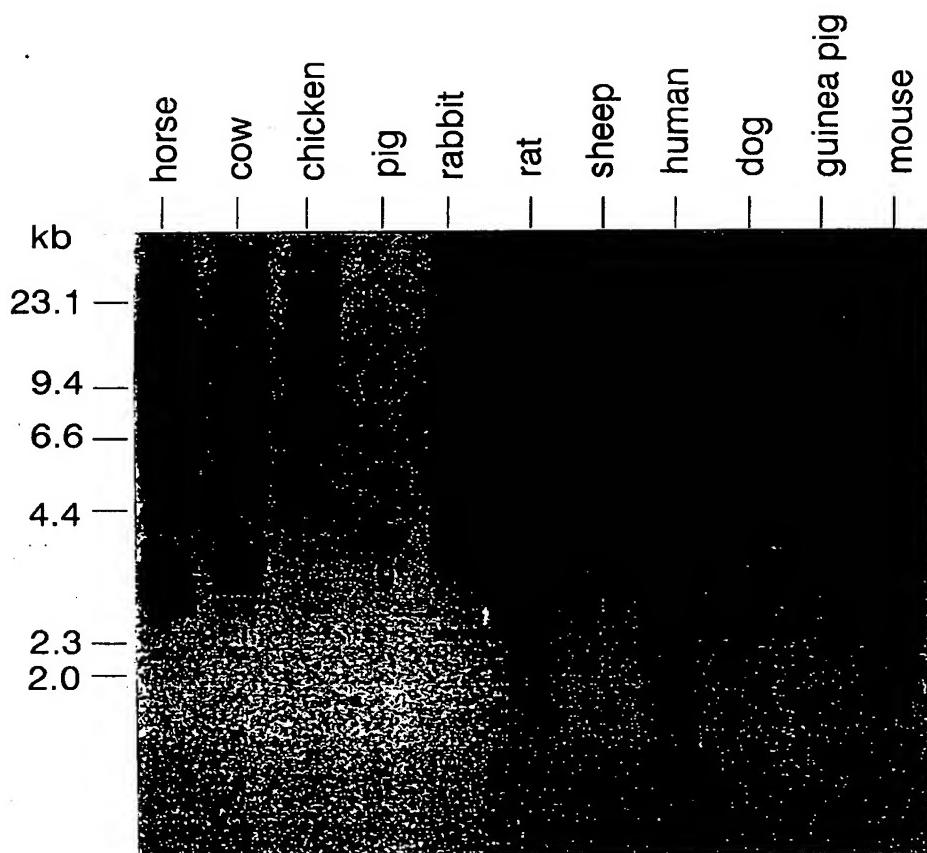


Figure 6



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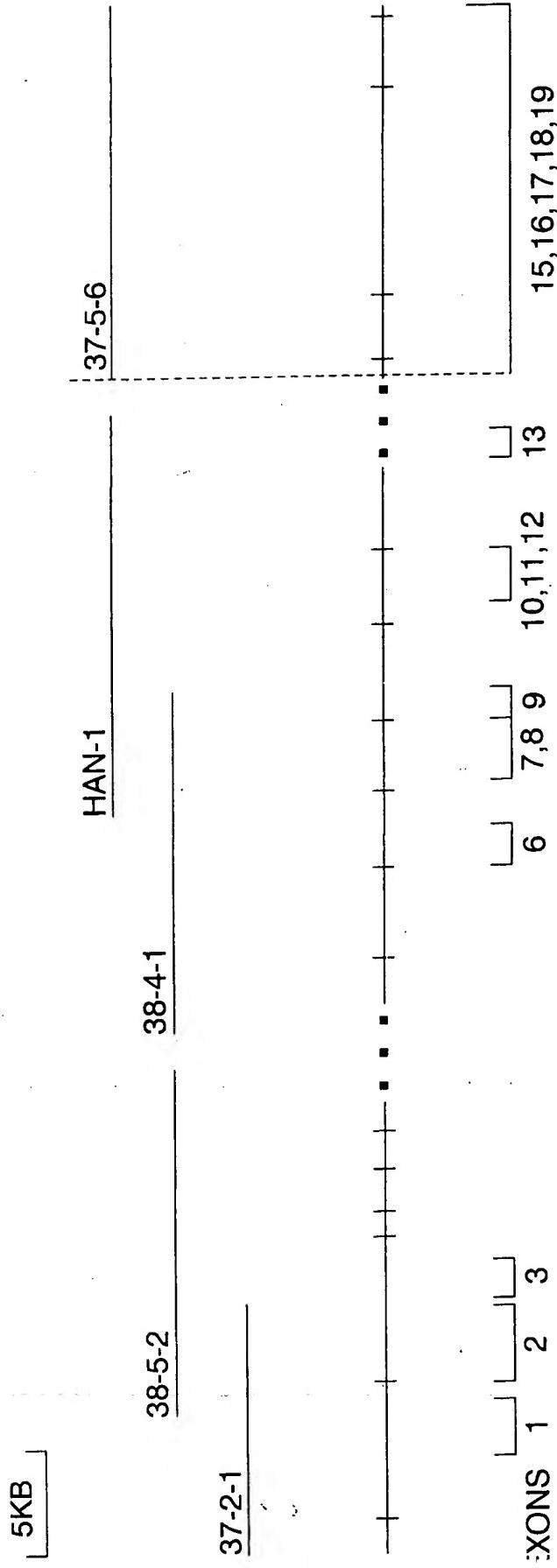


Cross-species conservation of the NR-2 gene

**Figure 7**



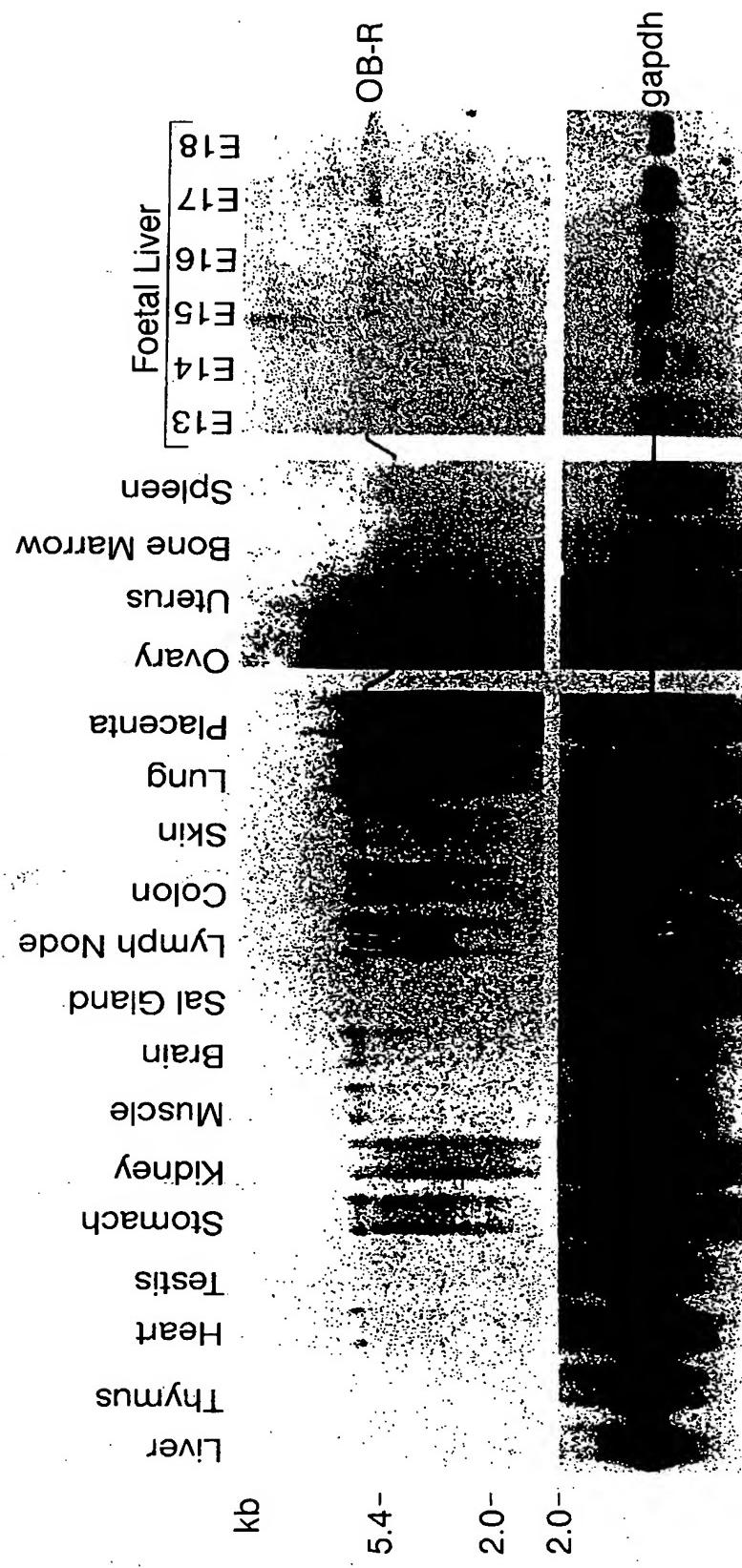
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**Figure 8**



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Expression of the Leptin Receptor (NR2) in murine tissues

## Figure 9

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